

FIG. 1

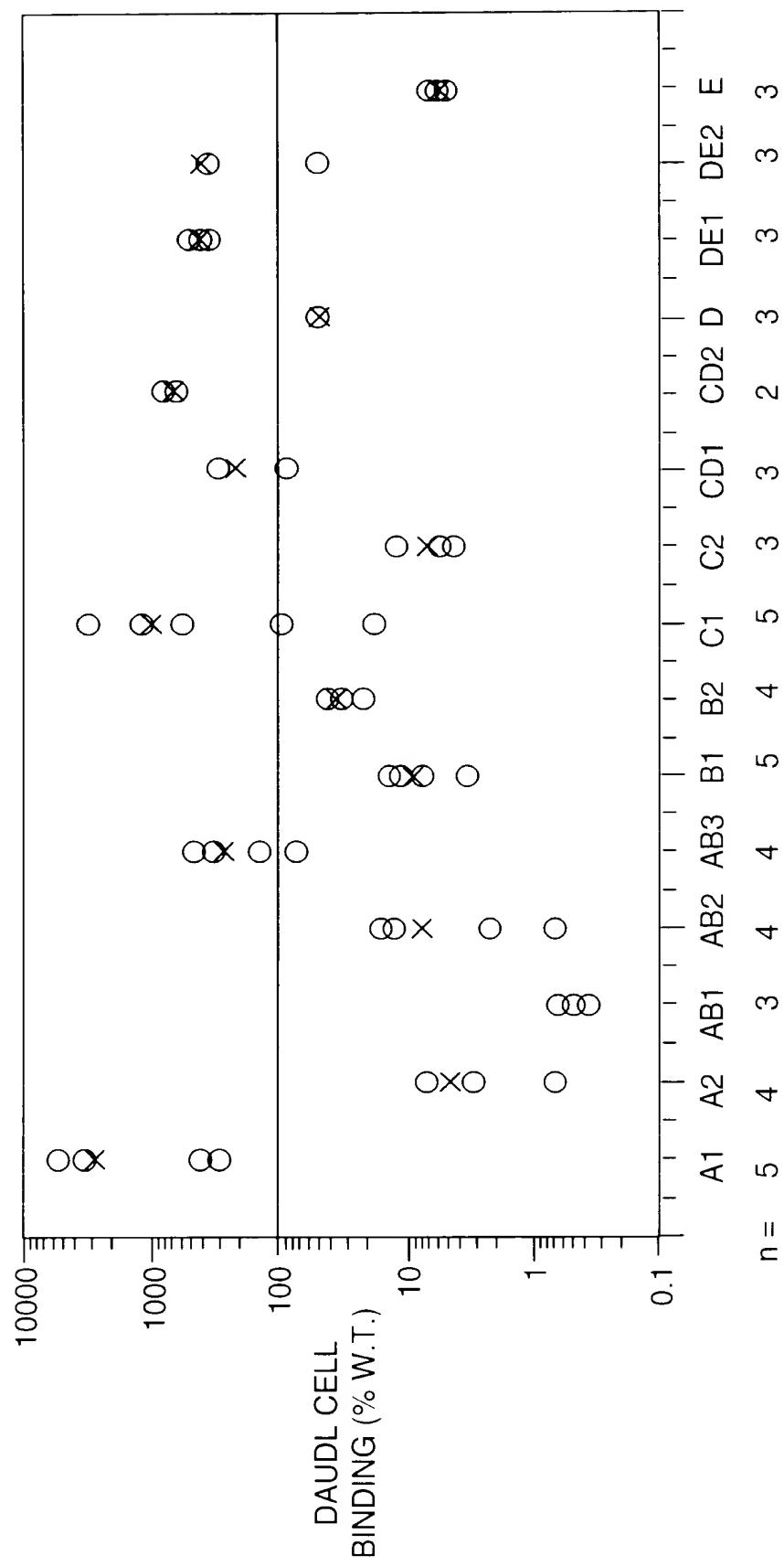


FIG. 2

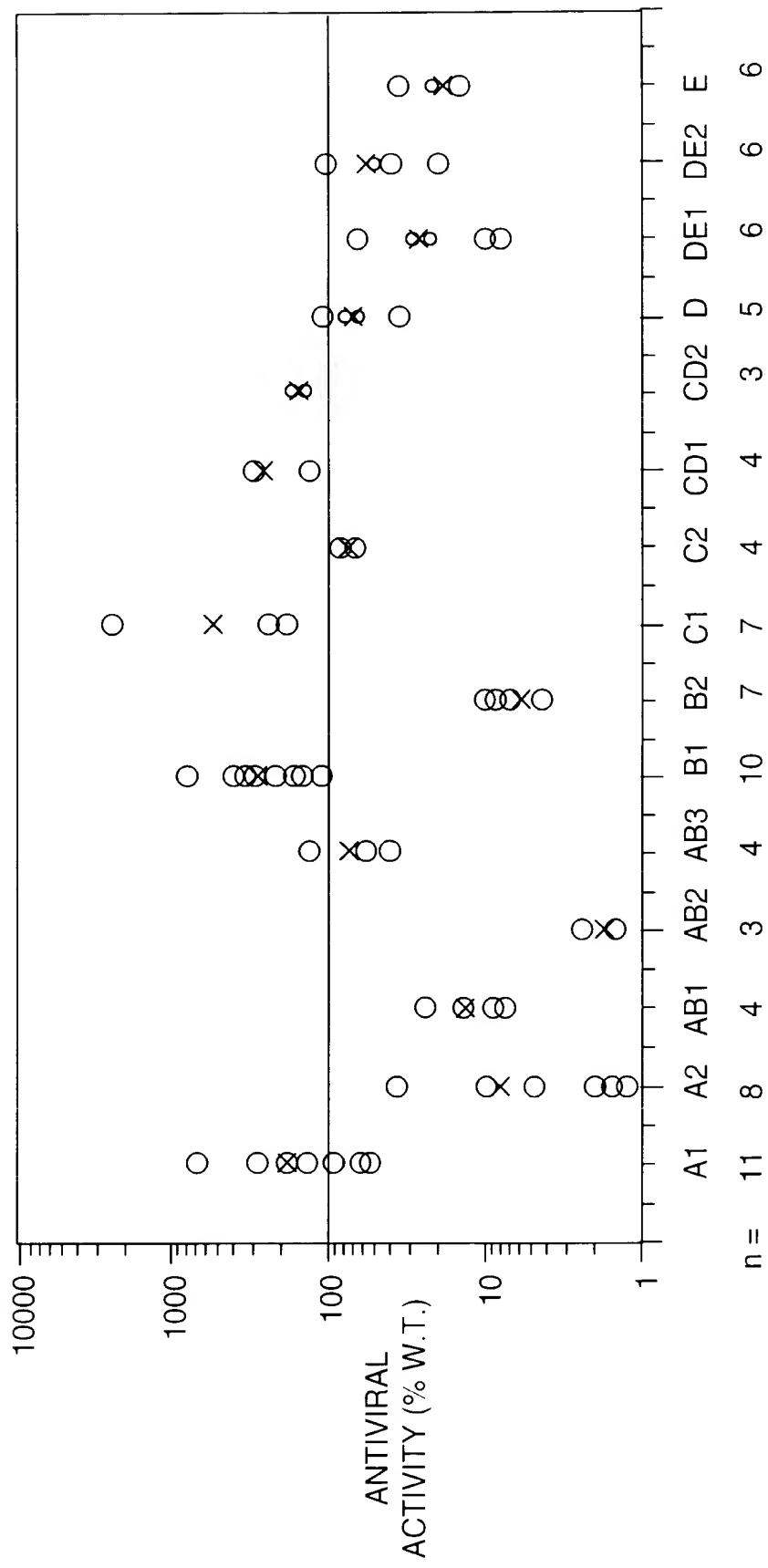


FIG. 3

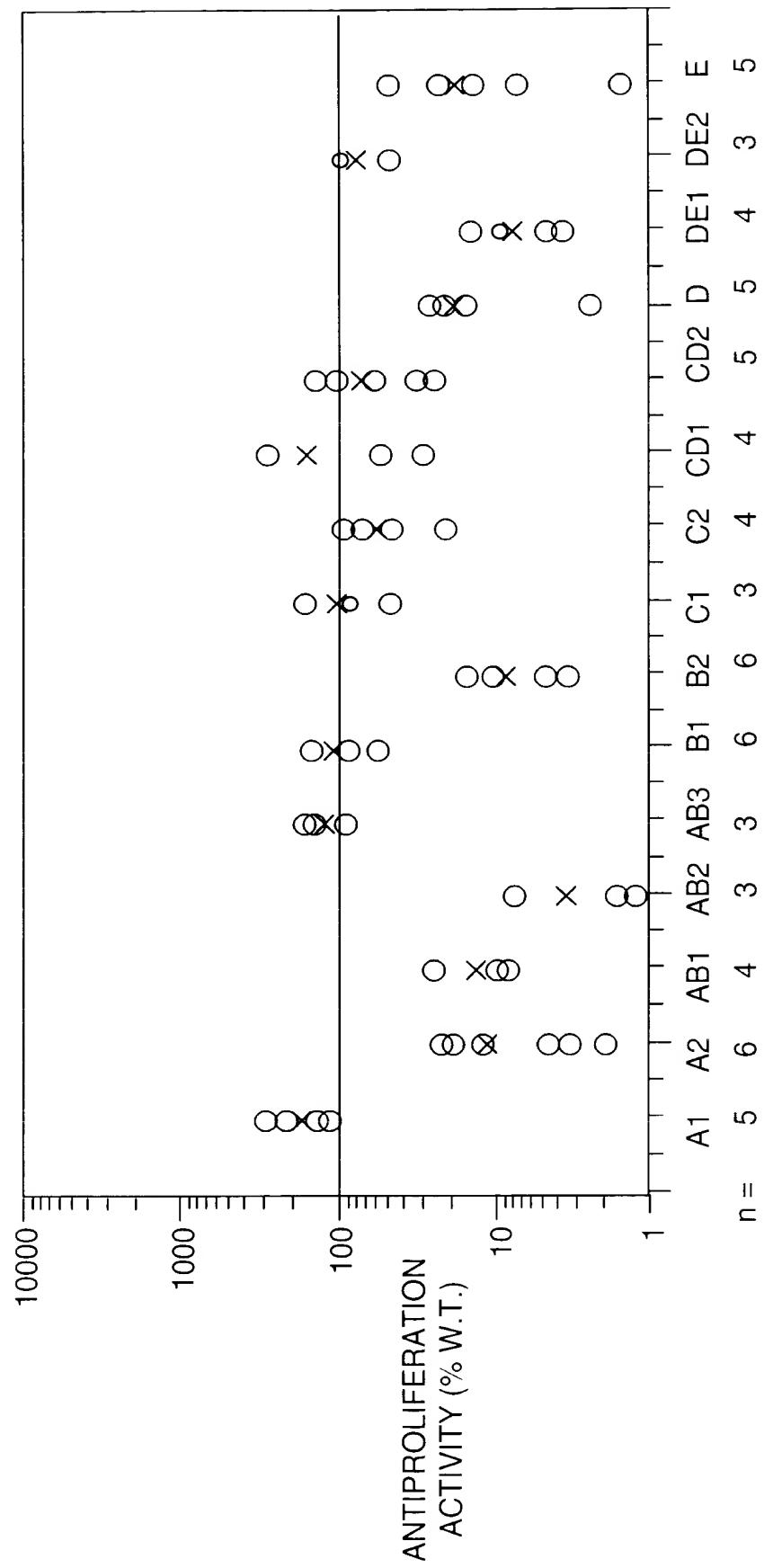


FIG. 4

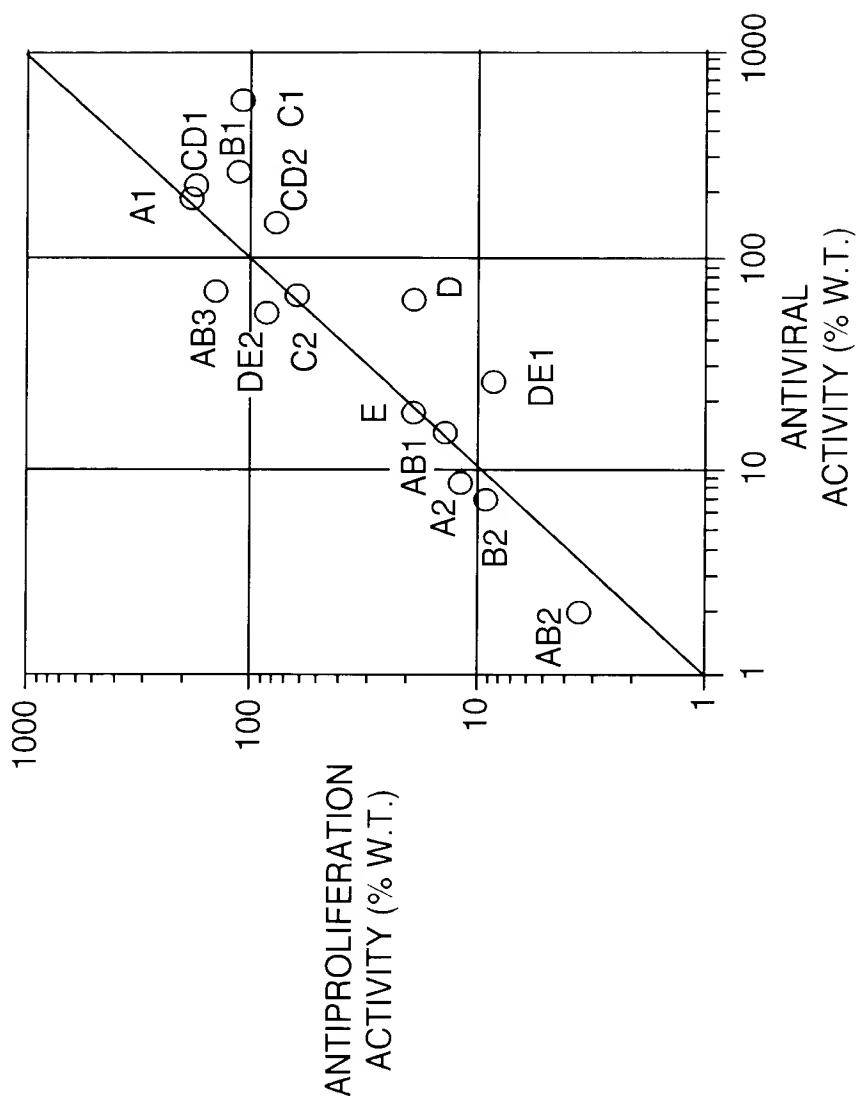


FIG. 5

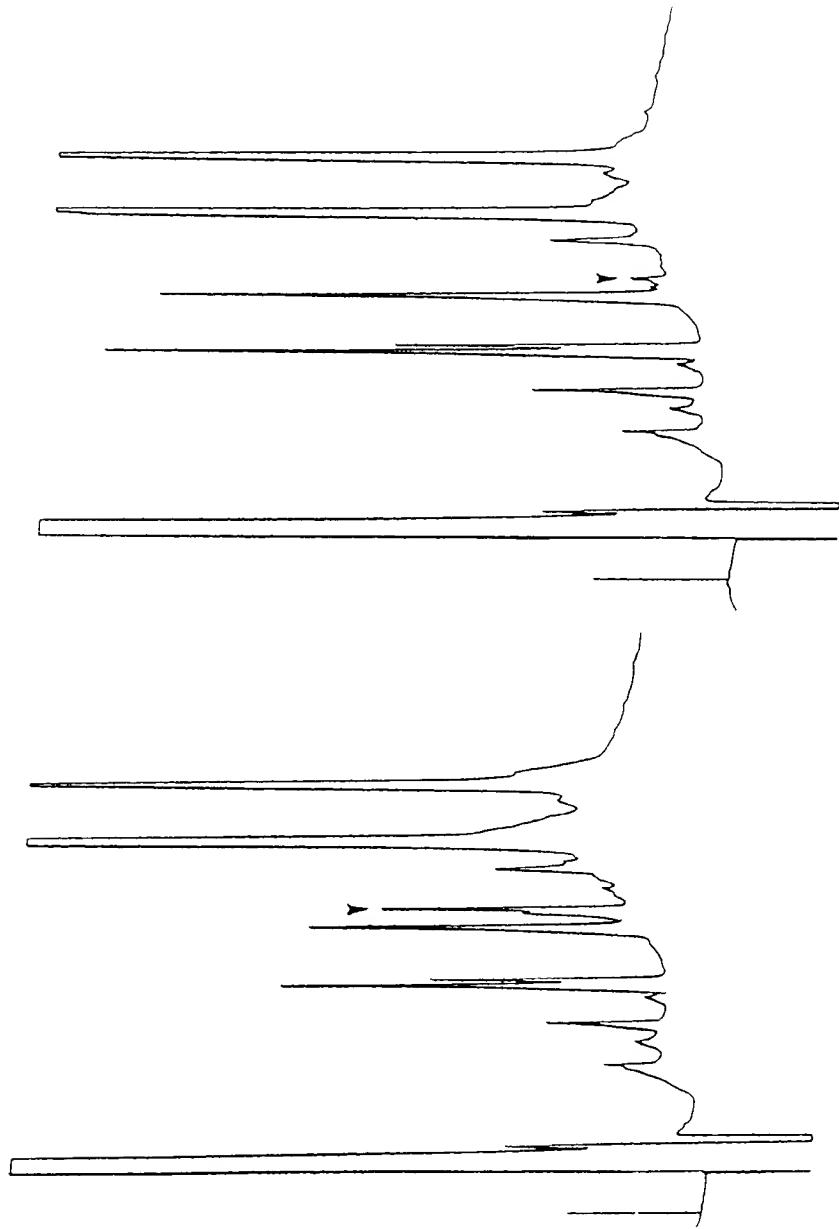


FIG. 6B

FIG. 6A

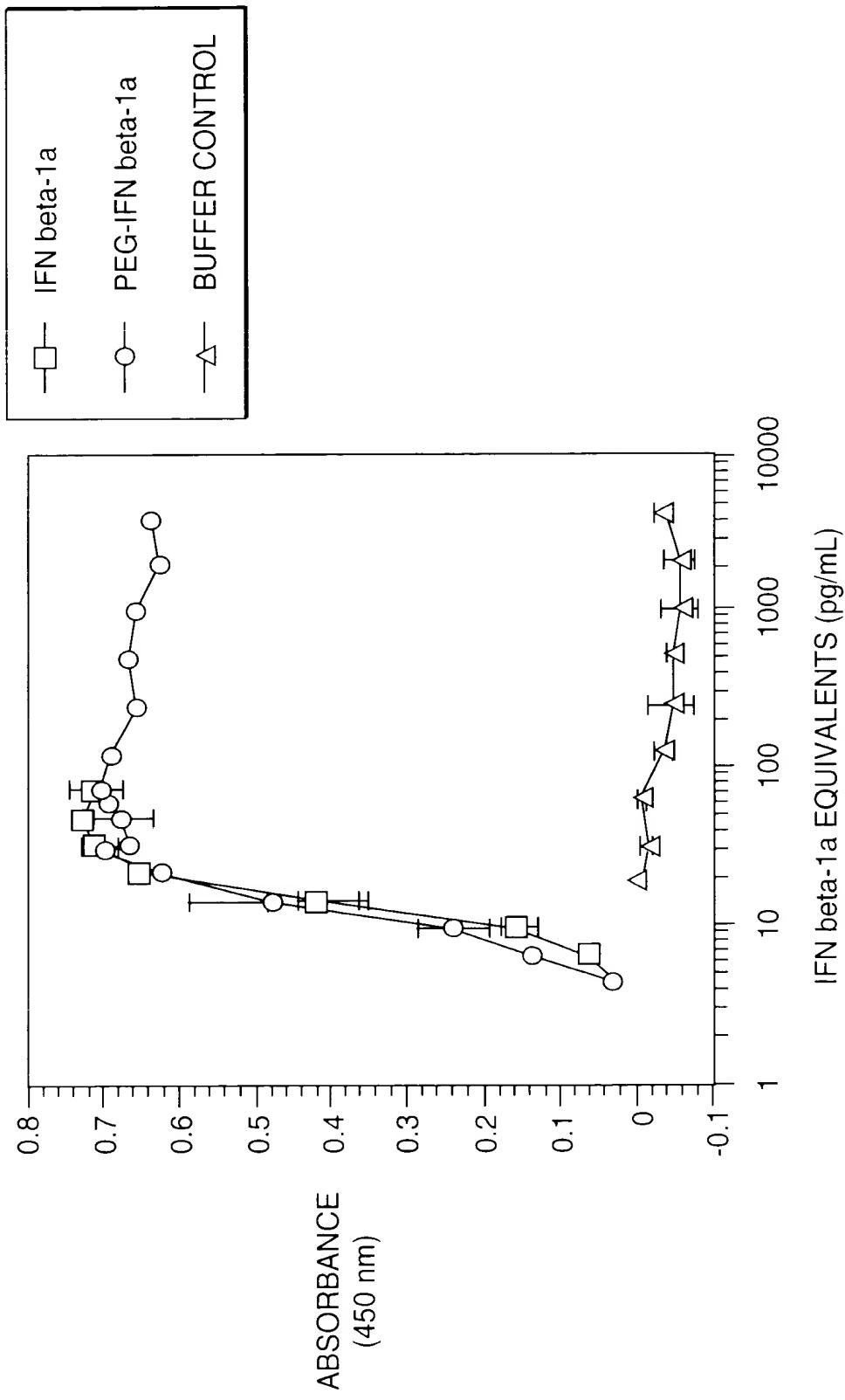


FIG. 7

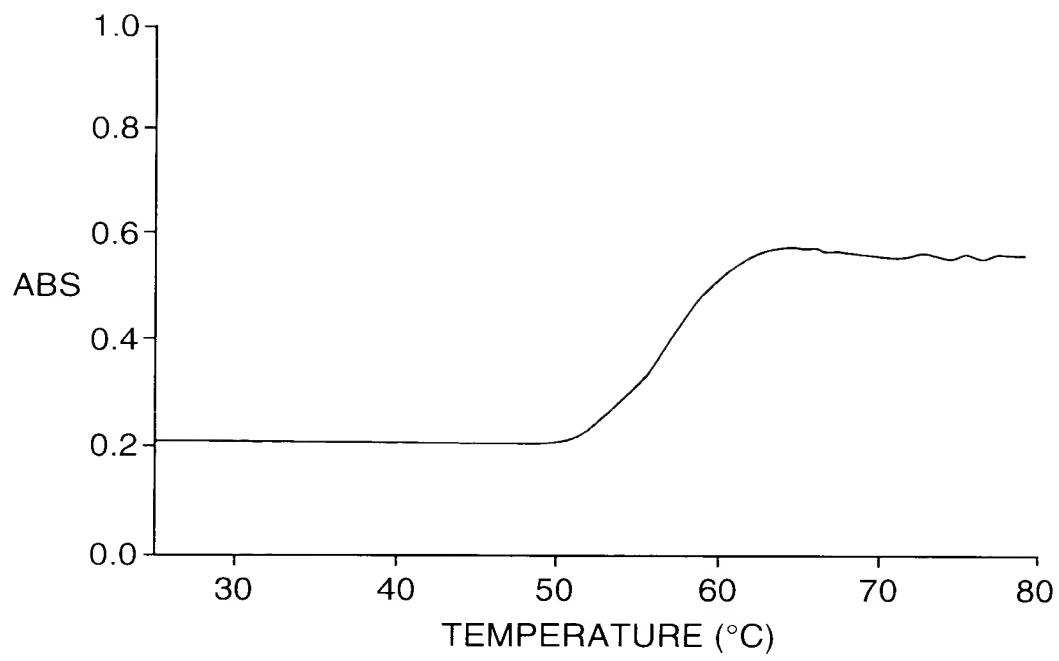


FIG. 8a

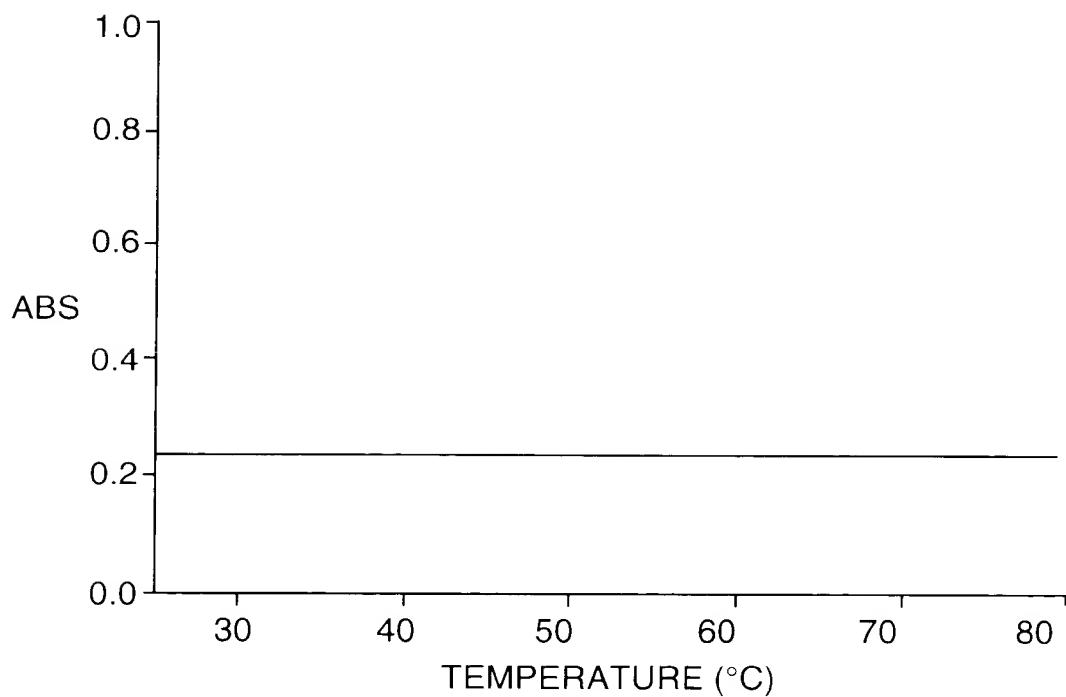


FIG. 8b

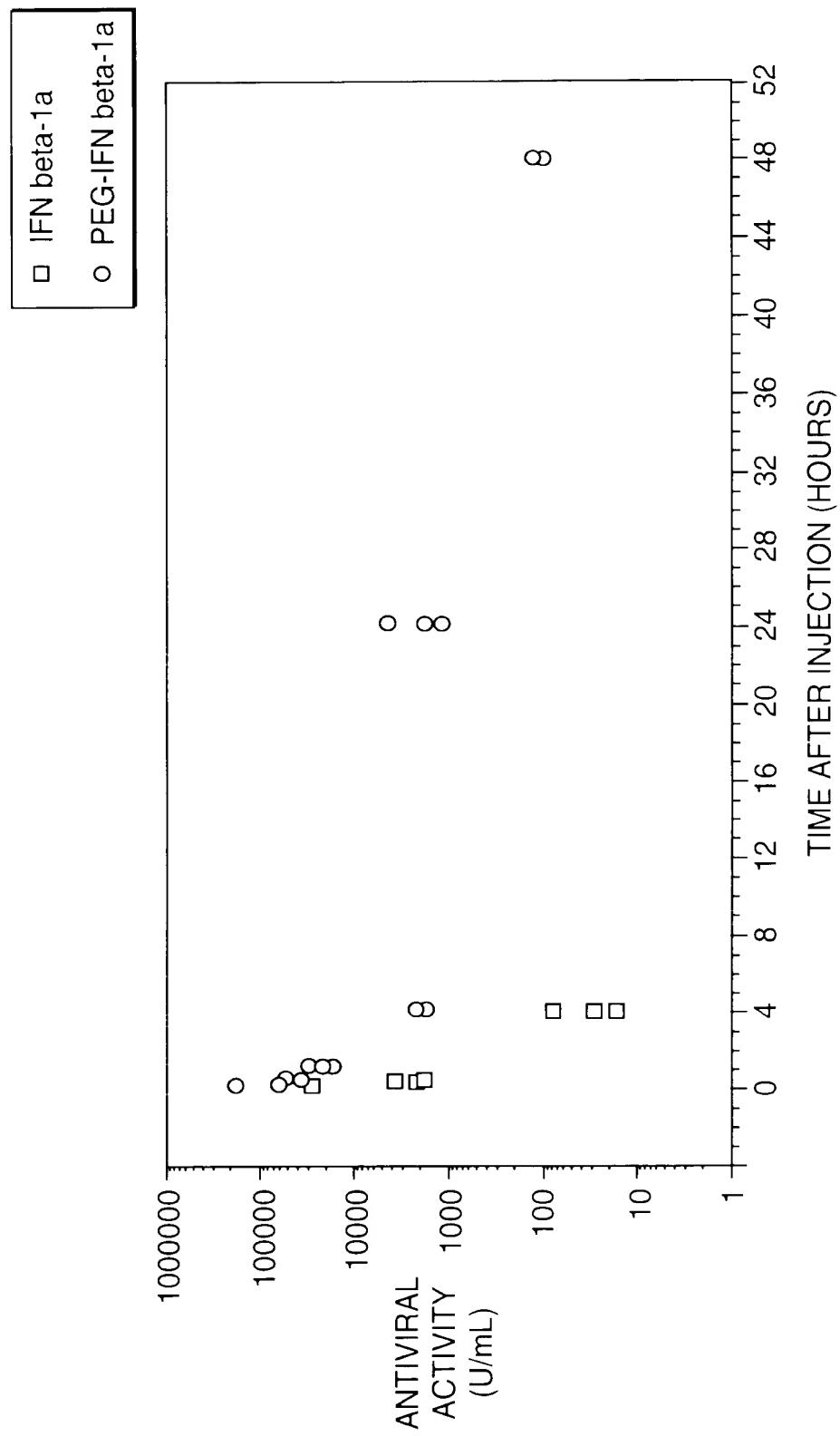


FIG. 9

1 TCCGGGGGCC ATCATCATCA TCATCATAGC TCCGGAGACG ATGATGACAA GATGAGCTAC
 AGGCCCCCGG TAGTAGTAGT AGTAGTATCG AGGCCTCTGC TACTACTGTT CTACTCGATG
 1►Ser GlyGlyH iHisHisHi sHisHisSer Ser GlyAspA spAspAspLy sMetSer Tyr

 61 AACTTGCTTG GATTCCCTACA AAGAACGAGC AATTTCACT GTCAGAAGCT CCTGTGGCAA
 TTGAACGAAC CTAAGGATGT TTCTTCGTCG TTAAAGTCA CAGTCTTCGA GGACACCGTT
 21►AsnLeuLeuG lypheLeuGI nArgSerSer AsnPheGI nC ysGlnLysLe uLeuTrpGI n

 121 TTGAATGGGA GGCTTGAATA CTGCCTCAAG GACAGGATGA ACTTTGACAT CCTGTGGAG
 AACTTACCCCT CCGAACTTAT GACGGAGTTC CTGTCCTACT TGAAACTGTA GGGACTCCCTC
 41►LeuAsnGlyA rglLeuGI uTy rCysLeuLys AspArgMetA snPheAspII eProGI uGI u

 181 ATTAAGCAGC TGACAGCTT CCAGAACGGAG GACGCCGCAT TGACCATCTA TGAGATGCTC
 TAATTGTCG ACAGTCGTCAA GGTCTTCCTC CTGCGGCGTA ACTGGTAGAT ACTCTACGAG
 61►IleLysGlnL euGI nGlnPh eGI nLysGlu AspAlaAlaL euThrIleTy rGI uMetLeu

 241 CAGAACATCT TTGCTATTT CAGACAAGAT TCATCTAGCA CTGGCTGGAA TGAGACTATT
 GTCTTGAGA AACGATAAAA GTCTGTTCTA AGTAGATCGT GACCGACCTT ACTCTGATAA
 81►GI nAsnIleP heAlaIlePh eArgGI nAsp SerSerSerT hrGlyTrpAs nGluThrIle

 301 GTTGAGAACCC TCCTGGCTAA TGTCTATCAT CAGATAAACCC ATCTGAAGAC AGTCTGGAA
 CAACTCTTGG AGGACCGATT ACAGATAGTA GTCTATTGG TAGACTTCTG TCAGGACCTT
 101►ValGluAsnL euLeuAlaAs nValTyrHis GlnIleAsnH iLeuLysTh rValLeuGI u

 361 GAAAAACTGG AGAAAGAAGA TTTCACCAAGG GGAAACTCA TGAGCAGTCT GCACCTGAAA
 CTTTTGACC TCTTTCTTCT AAAGTGGTCC CTTTTGAGT ACTCGTCAGA CGTGGACTTT
 121►GluLysLeuG lLysGluAs pPheThrArg GlyLysLeuM etSerSerLe uHisLeuLys

 421 AGATATTATG GGAGGATTCT GCATTACCTG AAGGCCAAGG AGTACAGTCA CTGTGCCTGG
 TCTATAATAC CCTCCTAAGA CGTAATGGAC TTCCGGTTCC TCATGTCAGT GACACGGACC
 141►ArgTyrTyrG lypheLeuIleLe uHisTyrLeu LysAlaLysG luTyrSerHi sCysAlaTrp

 481 ACCATAGTCA GAGTGGAAAT CCTAAGGAAC TTTTACTTCA TTAACAGACT TACAGGTTAC
 TGGTATCAGT CTCACCTTTA GGATTCCTTG AAAATGAAGT AATTGTCTGA ATGTCCAATG
 161►ThrIleValA rglValGluII eLeuArgAsn PheTyrPhel IleAsnArgLe uThrGlyTyr

 541 CTCCGAAAC
 GAGGCTTG
 181►LeuArgAsn

FIG. 10

